Bayesian modelling Bayesics

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Probability vs frequency

In frequentist statistic, "probability" is synonym for

long-term frequency under repeated sampling



What is probability?

Probability reflects incomplete information.

Quoting Finetti (1974)

Probabilistic reasoning — always to be understood as subjective — merely stems from our being uncertain about something.

Why opt for the Bayesian paradigm?

- Satisfies the likelihood principle
- Generative approach naturally extends to complex settings (hierarchical models)
- Uncertainty quantification and natural framework for prediction
- Capability to incorporate subject-matter expertise

Bayesian versus frequentist

Frequentist

- Parameters treated as fixed, data as random
 - true value of parameter θ is unknown.
- Target is point estimator

Bayesian

- Both parameters and data are random
 - inference is conditional on observed data
- Target is a distribution

Joint and marginal distribution

The joint density of data $oldsymbol{Y}$ and parameters $oldsymbol{ heta}$ is

$$p(\mathbf{Y}, \boldsymbol{\theta}) = p(\mathbf{Y} \mid \boldsymbol{\theta})p(\boldsymbol{\theta}) = p(\boldsymbol{\theta} \mid \mathbf{Y})p(\mathbf{Y})$$

where the marginal $p(\boldsymbol{Y}) = \int_{\boldsymbol{\Theta}} p(\boldsymbol{Y}, \boldsymbol{\theta}) \mathrm{d} \boldsymbol{\theta}$.

Posterior

Using Bayes' theorem, the posterior density is

$$p(\boldsymbol{\theta} \mid \boldsymbol{Y}) = \frac{p(\boldsymbol{Y} \mid \boldsymbol{\theta}) \times p(\boldsymbol{\theta})}{\int p(\boldsymbol{Y} \mid \boldsymbol{\theta}) p(\boldsymbol{\theta}) d\boldsymbol{\theta}},$$

meaning that

posterior
$$\propto$$
 likelihood \times prior

Evaluating the marginal likelihood p(Y), is challenging when θ is high-dimensional.

Updating beliefs and sequentiality

By Bayes' rule, we can consider *updating* the posterior by adding terms to the likelihood, noting that for independent y_1 and y_2 ,

$$p(oldsymbol{ heta} \mid oldsymbol{y}_1, oldsymbol{y}_2) \propto p(oldsymbol{y}_2 \mid oldsymbol{ heta}) p(oldsymbol{ heta} \mid oldsymbol{y}_1)$$

The posterior is be updated in light of new information.

Binomial distribution

A binomial variable with probability of success $heta \in [0,1]$ has mass function

$$f(y; heta) = inom{n}{y} heta^y (1- heta)^{n-y}, \qquad y=0,\dots,n.$$

Moments of the number of successes out of n trials are

$$\mathsf{E}(Y \mid \theta) = n\theta, \quad \mathsf{Va}(Y \mid \theta) = n\theta(1-\theta).$$

The binomial coefficient $\binom{n}{y}=n!/\{(n-y)!y!\}$, where $n!=\Gamma(n+1)$.

Beta distribution

The beta distribution with shapes $\alpha > 0$ and $\beta > 0$, denoted beta (α, β) , has density

$$f(y) = rac{\Gamma(lpha+eta)}{\Gamma(lpha)\Gamma(eta)} y^{lpha-1} (1-y)^{eta-1}, \qquad y \in [0,1]$$

- expectation: $\alpha/(\alpha+\beta)$;
- mode $(\alpha-1)/(\alpha+\beta-2)$ if $\alpha,\beta>1$, else, 0,1 or none;
- variance: $\alpha\beta/\{(\alpha+\beta)^2(\alpha+\beta+1)\}.$

Beta-binomial example

We write $Y \sim \mathsf{binom}(n, \theta)$ for $\theta \in [0, 1]$; the likelihood is

$$L(heta;y) = inom{n}{y} heta^y (1- heta)^{n-y}.$$

Consider a beta prior, $\theta \sim \mathsf{beta}(\alpha, \beta)$, with density

$$p(heta) = rac{\Gamma(lpha+eta)}{\Gamma(lpha)\Gamma(eta)} heta^{lpha-1}(1- heta)^{eta-1}.$$

Density versus likelihood

The binomial distribution is discrete with support $0, \ldots, n$, whereas the likelihood is continuous over $\theta \in [0, 1]$.

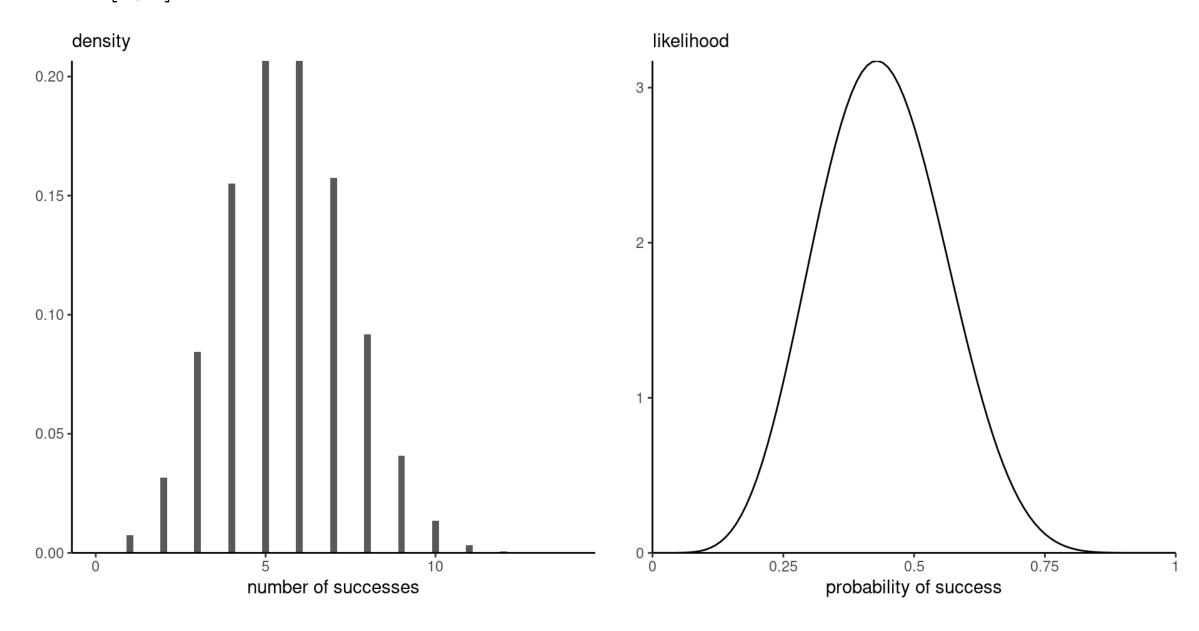


Figure 1: Binomial density function (left) and scaled likelihood function (right).

If the density or mass function integrates to 1 over the range of Y, the integral of the likelihood over θ does not.

Posterior density and proportionality

Any term not a function of θ can be dropped, since it will absorbed by the normalizing constant. The posterior density is proportional to

$$egin{aligned} L(heta;y)p(heta) & \stackrel{ heta}{\propto} heta^y (1- heta)^{n-y} imes heta^{lpha-1} (1- heta)^{eta-1} \ &= heta^{y+lpha-1} (1- heta)^{n-y+eta-1} \end{aligned}$$

the kernel of a beta density with shape parameters $y+\alpha$ and $n-y+\beta$.

The symbol ∞ , for proportionality, means dropping all terms not an argument of the left hand side.

Marginal likelihood

The marginal likelihood for the $Y \mid P = p \sim \mathsf{binom}(n,p)$ model with prior $P \sim \mathsf{beta}(\alpha,\beta)$ is

$$p_Y(y) = inom{n}{y} rac{\mathrm{beta}(lpha+y,eta+n-y)}{\mathrm{beta}(lpha,eta)}, \quad y \in \{0,\dots,n\}.$$

where $beta(\alpha, \beta) = \Gamma(\alpha)\Gamma(\beta)/\Gamma(\alpha+\beta)$ is the beta function.

Experiments and likelihoods

Consider the following sampling mechanism, which lead to k successes out of n independent trials, with the same probability of success θ .

- 1. Bernoulli: sample fixed number of observations with $L(\theta;y)= heta^k(1- heta)^{n-k}$
- 2. binomial: same, but record only total number of successes so $L(\theta;y)=\binom{n}{k}\theta^k(1-\theta)^{n-k}$
- 3. negative binomial: sample data until you obtain a predetermined number of successes, whence $L(\theta;y)=\binom{n-1}{k-1}\theta^k(1-\theta)^{n-k}$

Likelihood principle

Two likelihoods that are proportional, up to a constant not depending on unknown parameters, yield the same evidence.

In all cases, $L(\theta;y) \overset{\theta}{\propto} \theta^k (1-\theta)^{n-k}$, so these yield the same inference for Bayesian.

For a more in-depth discussion, see Section 6.3.2 of Casella & Berger (2002)

Integration

We could approximate the marginal likelihood through either

- numerical integration (cubature)
- Monte Carlo simulations

In more complicated models, we will try to sample observations by bypassing completely this calculation.

The likelihood terms can be small (always less than one and decreasing for discrete data), so watch out for numerical overflow when evaluating normalizing constants.

Numerical example of (Monte Carlo) integration

```
[1] 1.066906e-05
```

2 beta(y + alpha, n - y + beta)

```
1 # Monte Carlo integration
2 mean(unnormalized_posterior(runif(1e5)))
```

[1] 1.061693e-05

Marginal posterior

In multi-parameter models, additional integration is needed to get the marginal posterior

$$p(heta_j \mid oldsymbol{y}) = \int p(oldsymbol{ heta} \mid oldsymbol{y}) \mathrm{d}oldsymbol{ heta}_{-j}.$$

Marginalization is trivial when we have a joint sample: simply keep the column corresponding to θ_j .

Prior, likelihood and posterior

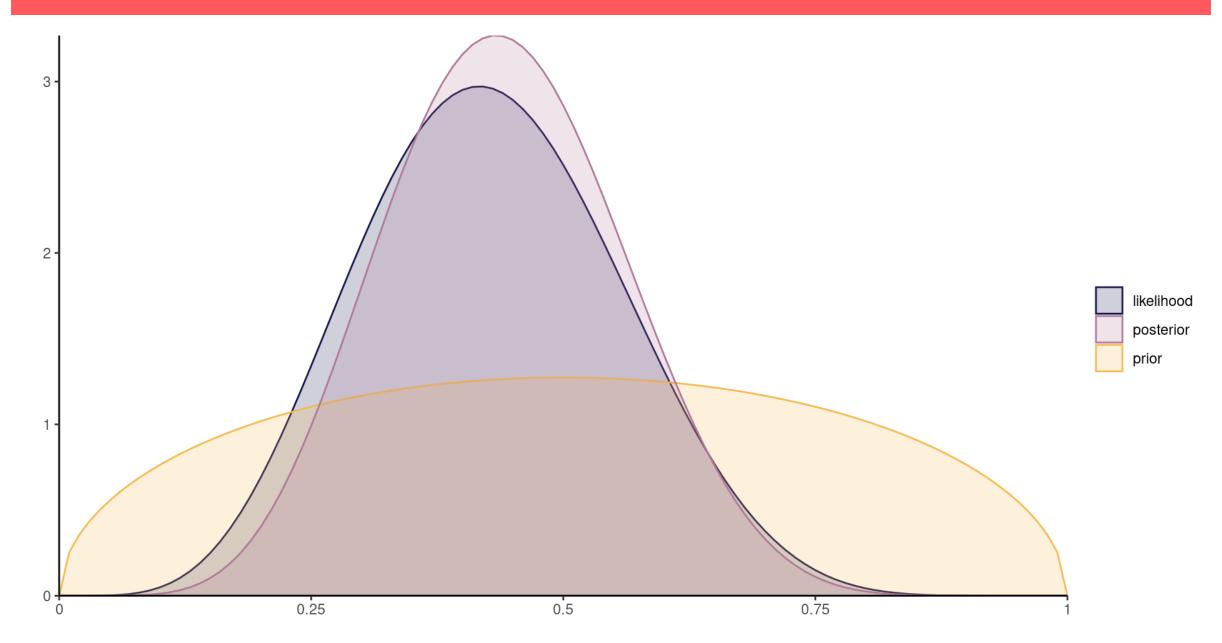


Figure 2: Scaled Binomial likelihood for six successes out of 14 trials, beta(3/2,3/2) prior and corresponding posterior distribution from a beta-binomial model.

Proper prior

We could define the posterior simply as the normalized product of the likelihood and some prior function.

The prior function need not even be proportional to a density function (i.e., integrable as a function of θ).

For example,

- $p(\theta) \propto \theta^{-1} (1-\theta)^{-1}$ is improper because it is not integrable.
- $p(\theta) \propto 1$ is a proper prior over [0,1] (uniform).

Validity of the posterior

- ullet The marginal likelihood does not depend on $oldsymbol{ heta}$
 - (a normalizing constant)
- For the posterior density to be proper,
 - the marginal likelihood must be a finite!
 - in continuous models, the posterior is proper whenever the prior function is proper.

Different priors give different posteriors

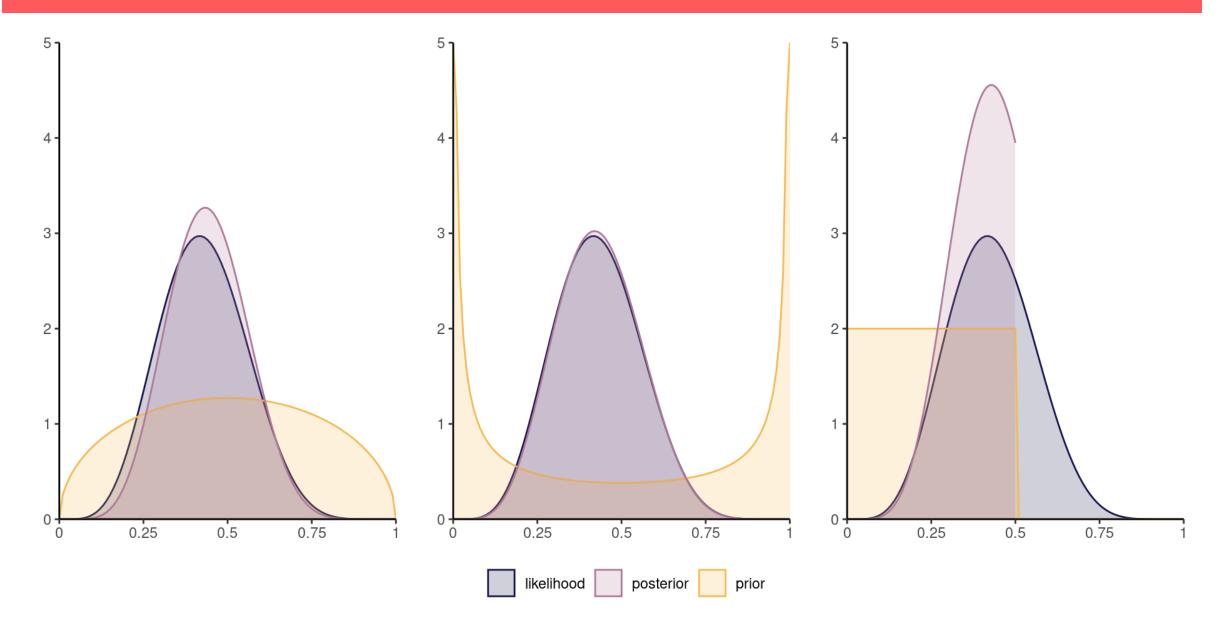


Figure 3: Scaled binomial likelihood for six successes out of 14 trials, with beta(3/2, 3/2) (left), beta(1/4, 1/4) (middle) and unif[0, 1/2] (right) priors and posterior density.

Role of the prior

The posterior is beta, with expected value

$$\mathsf{E}(heta \mid y) = wrac{y}{n} + (1-w)rac{lpha}{lpha + eta}, \ w = rac{n}{n+lpha + eta}$$

a weighted average of

- the maximum likelihood estimator and
- the prior mean.

Posterior concentration

Except for stubborn priors, the likelihood contribution dominates in large samples. The impact of the prior is then often negligible.

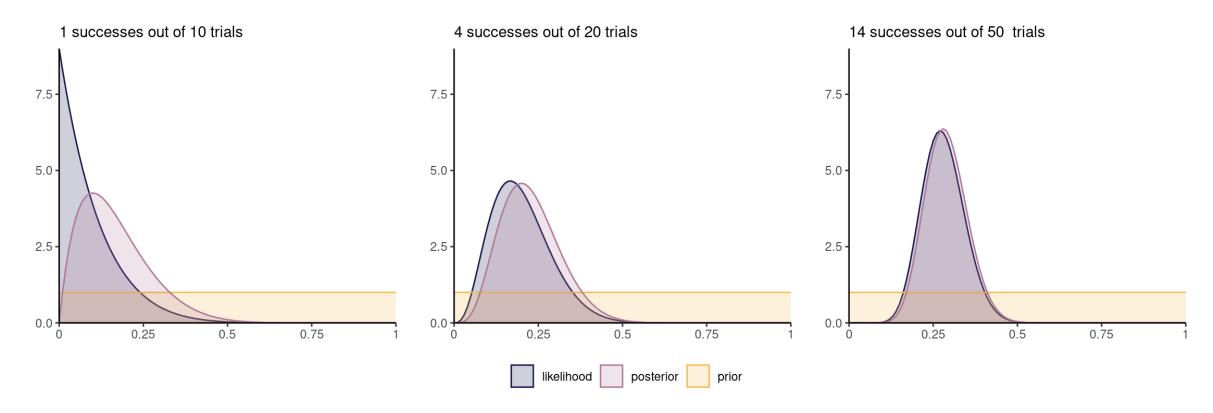


Figure 4: Beta posterior and binomial likelihood with a uniform prior for increasing number of observations (from left to right).

Model comparison

Suppose that we have models \mathcal{M}_m $(m=1,\ldots,M)$ to be compared, with parameter vectors $\boldsymbol{\theta}^{(m)}$ and data vector \boldsymbol{y} and prior probability $\Pr(\mathcal{M}_m)$.

The posterior odds for models \mathcal{M}_i vs \mathcal{M}_j is

$$rac{\Pr(\mathcal{M}_i \mid oldsymbol{y})}{\Pr(\mathcal{M}_j \mid oldsymbol{y})} = rac{p(oldsymbol{y} \mid \mathcal{M}_i)}{p(oldsymbol{y} \mid \mathcal{M}_j)} rac{\Pr(\mathcal{M}_i)}{\Pr(\mathcal{M}_j)}$$

equal to the Bayes factor BF_{ij} times the prior odds.

Bayes factors

The Bayes factor is the ratio of marginal likelihoods, as

$$p(oldsymbol{y} \mid \mathcal{M}_i) = \int p(y \mid oldsymbol{ heta}^{(i)}, \mathcal{M}_i) p(oldsymbol{ heta}^{(i)} \mid \mathcal{M}_i) \mathrm{d}oldsymbol{ heta}^{(i)}.$$

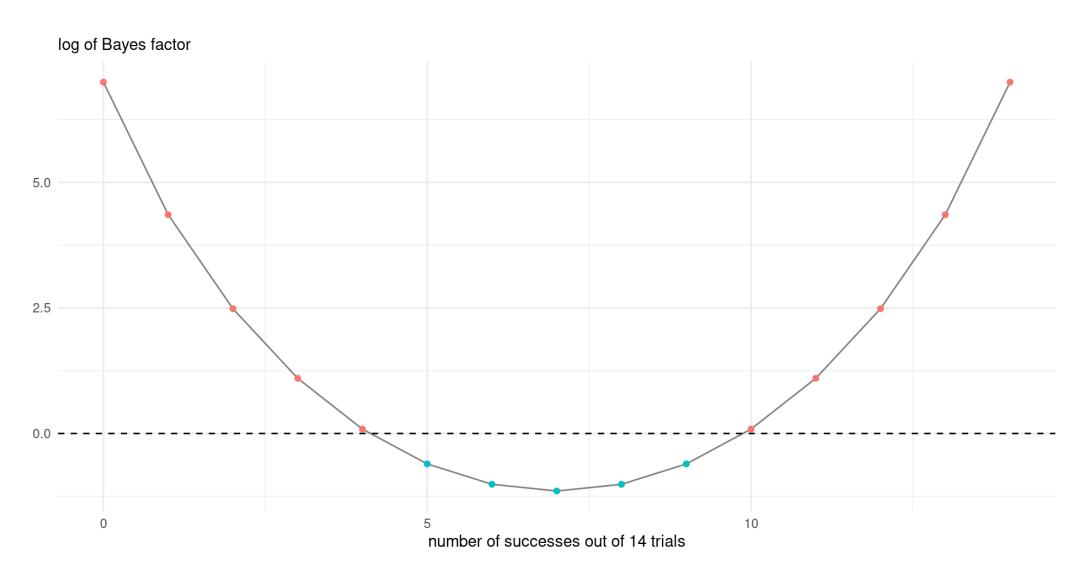
Values of $\mathsf{BF}_{ij} > 1$ correspond to model \mathcal{M}_i being more likely than \mathcal{M}_i .

- Strong dependence on the prior $p(\boldsymbol{\theta}^{(i)} \mid \mathcal{M}_i)$.
- Must use proper priors.

Bayes factor for the binomial model

Consider two models with $Y \mid P^{(i)} = p \sim \mathsf{binom}(n,p)$ and

- ullet $P^{(1)}\sim \mathsf{unif}(0,1)$
- ullet $P^{(2)}\sim 1_{p=0.5}.$



Summarizing posterior distributions

The output of the Bayesian learning will be either of:

- 1. a fully characterized distribution (in toy examples).
- 2. a numerical approximation to the posterior distribution.
- 3. an exact or approximate sample drawn from the posterior distribution.

Bayesian inference in practice

Most of the field revolves around the creation of algorithms that either

- circumvent the calculation of the normalizing constant
 - (Monte Carlo and Markov chain Monte Carlo methods)
- provide accurate numerical approximation, including for marginalizing out all but one parameter.
 - (integrated nested Laplace approximations, variational inference, etc.)

Predictive distributions

Define the posterior predictive,

$$p(y_{ ext{new}} \mid oldsymbol{y}) = \int_{oldsymbol{\Theta}} p(y_{ ext{new}} \mid oldsymbol{ heta}) p(oldsymbol{ heta} \mid oldsymbol{y}) \mathrm{d}oldsymbol{ heta}$$

and the prior predictive

$$p(y_{
m new}) = \int_{oldsymbol{\Theta}} p(y_{
m new} \mid oldsymbol{ heta}) p(oldsymbol{ heta}) {
m d}oldsymbol{ heta}$$

is useful for determining whether the prior is sensical.

Analytical derivation of predictive distribution

Given the beta(a,b) prior or posterior, the predictive for n_{new} trials is beta-binomial with density

$$egin{aligned} p(y_{ ext{new}} \mid y) &= \int_0^1 inom{n_{ ext{new}}}{y_{ ext{new}}} rac{ heta^{a+y_{ ext{new}}-1}(1- heta)^{b+n_{ ext{new}}-y_{ ext{new}}-1}}{ ext{Be}(a,b)} \mathrm{d} heta \ &= inom{n_{ ext{new}}}{y_{ ext{new}}} rac{ ext{Be}(a+y_{ ext{new}},b+n_{ ext{new}}-y_{ ext{new}})}{ ext{Be}(a,b)} \end{aligned}$$

Replace $a=y+\alpha$ and $b=n-y+\beta$ to get the posterior predictive distribution.

Posterior predictive distribution

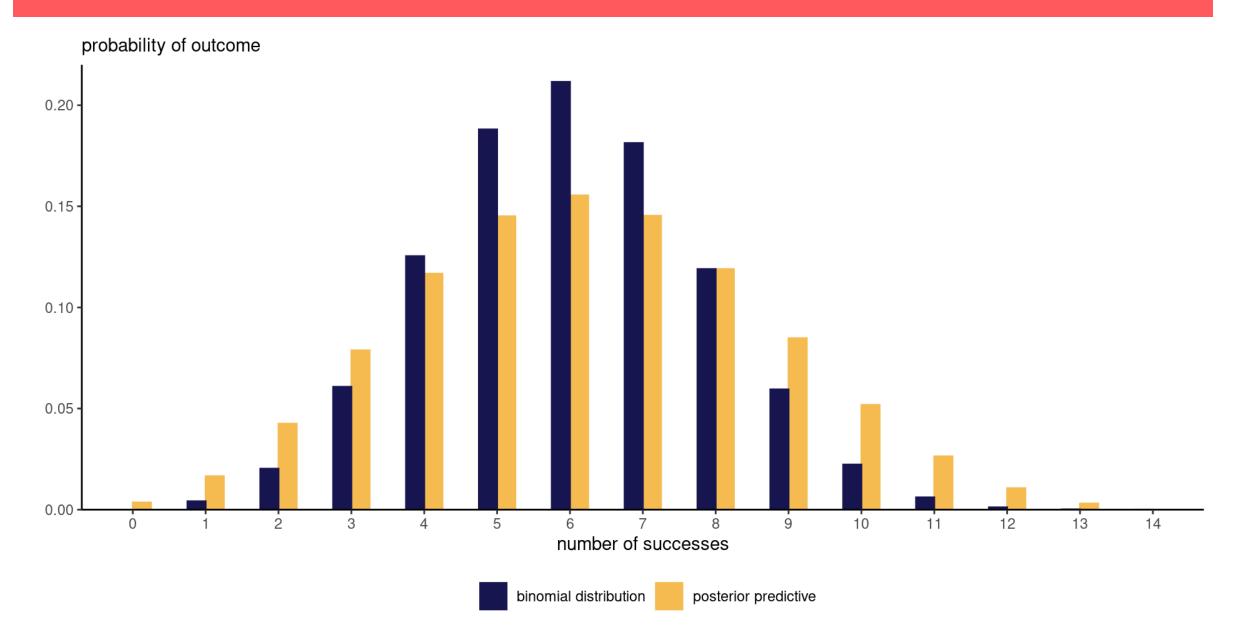


Figure 5: Beta-binomial posterior predictive distribution with corresponding binomial mass function evaluated at the maximum likelihood estimator.

Posterior predictive distribution via simulation

The posterior predictive carries over the parameter uncertainty so will typically be wider and overdispersed relative to the corresponding distribution.

Given a draw θ^* from the posterior, simulate a new observation from the distribution $f(y_{\text{new}}; \theta^*)$.

```
1 npost <- 1e4L
2 # Sample draws from the posterior distribution
3 post_samp <- rbeta(n = npost, y + alpha, n - y + beta)
4 # For each draw, sample new observation
5 post_pred <- rbinom(n = npost, size = n, prob = post_samp)</pre>
```

The beta-binomial is used to model overdispersion in binary regression models.

Summarizing posterior distributions

The output of a Bayesian procedure is a distribution for the parameters given the data.

We may wish to return different numerical summaries (expected value, variance, mode, quantiles, ...)

The question: which point estimator to return?

Decision theory and loss functions

A loss function $c(\theta, v) : \Theta \mapsto \mathbb{R}^k$ assigns a weight to each value θ , corresponding to the regret or loss.

The point estimator $\widehat{m{v}}$ is the minimizer of the expected loss

$$egin{aligned} \widehat{oldsymbol{v}} &= \underset{oldsymbol{v}}{\operatorname{argmin}} \mathsf{E}_{oldsymbol{\Theta} \mid oldsymbol{Y}} \{ c(oldsymbol{ heta}, oldsymbol{v}) \} \ &= \underset{oldsymbol{v}}{\operatorname{argmin}} \int_{oldsymbol{\Theta}} c(oldsymbol{ heta}, oldsymbol{v}) p(oldsymbol{ heta} \mid oldsymbol{y}) \mathrm{d}oldsymbol{ heta} \end{aligned}$$

Point estimators and loss functions

In a univariate setting, the most widely used point estimators are

- mean: quadratic loss $c(\theta, v) = (\theta v)^2$
- ullet median: absolute loss c(heta, v) = | heta v|
- mode: 0-1 loss $c(\theta, v) = 1 \mathrm{I}(v = \theta)$

The posterior mode $m{ heta}_{
m map} = {
m argmax}_{m{ heta}} p(m{ heta} \mid m{y})$ is the maximum a posteriori or MAP estimator.

Measures of central tendency

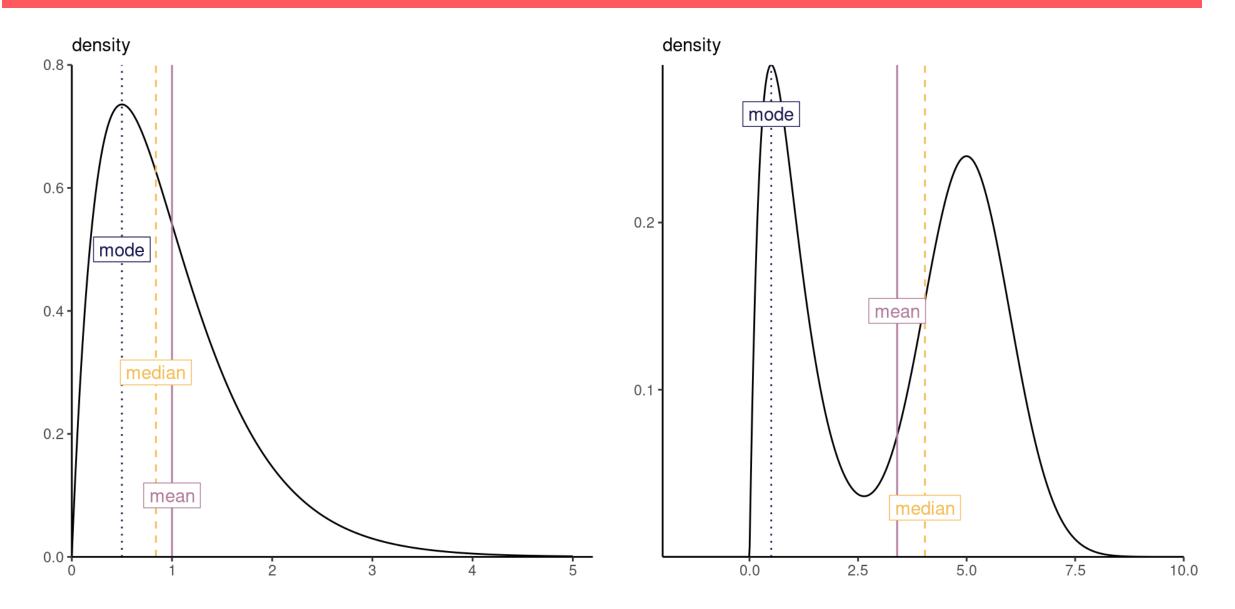


Figure 6: Point estimators from a right-skewed distribution (left) and from a multimodal distribution (right).

Example of loss functions

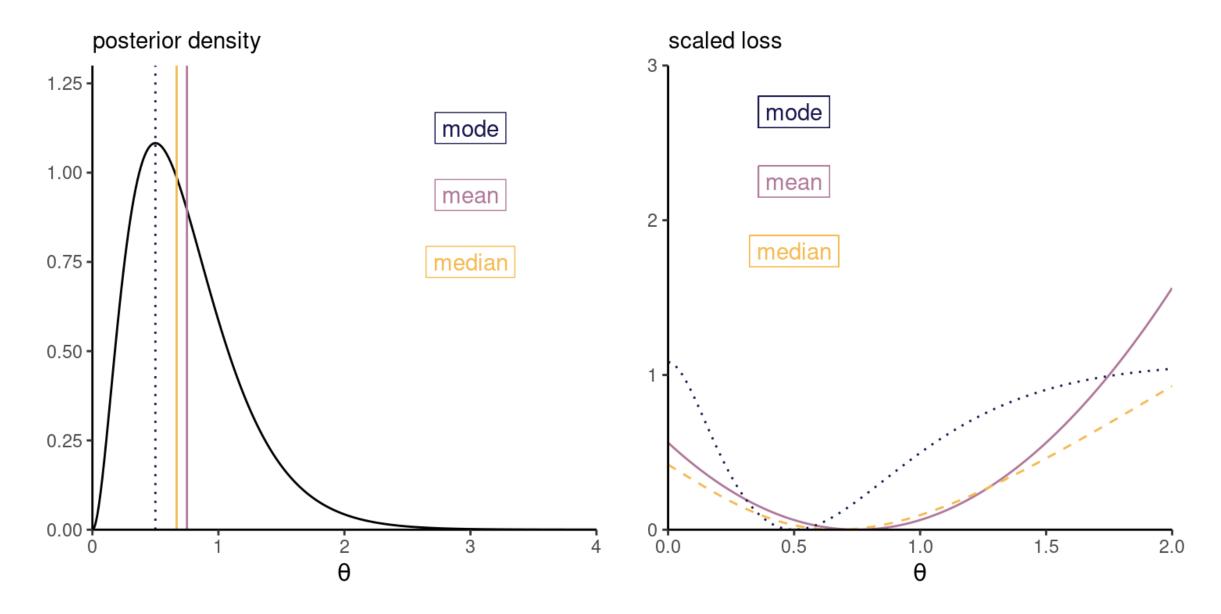


Figure 7: Posterior density with mean, mode and median point estimators (left) and corresponding loss functions, scaled to have minimum value of zero (right).

Credible regions

The freshman dream comes true!

A $1-\alpha$ credible region give a set of parameter values which contains the "true value" of the parameter $m{ heta}$ with probability $1-\alpha$.

Caveat: McElreath (2020) suggests the term 'compatibility', as it

returns the range of parameter values compatible with the model and data.

Which credible intervals?

Multiple $1-\alpha$ intervals, most common are

- ullet equitailed: region lpha/2 and 1-lpha/2 quantiles and
- highest posterior density interval (HPDI), which gives the smallest interval $(1-\alpha)$ probability

If we accept to have more than a single interval, the highest posterior density region can be a set of disjoint intervals. The HDPI is more sensitive to the number of draws and more computationally intensive (see **R** package HDinterval). See Hyndman (1996) for computations.

Illustration of credible regions

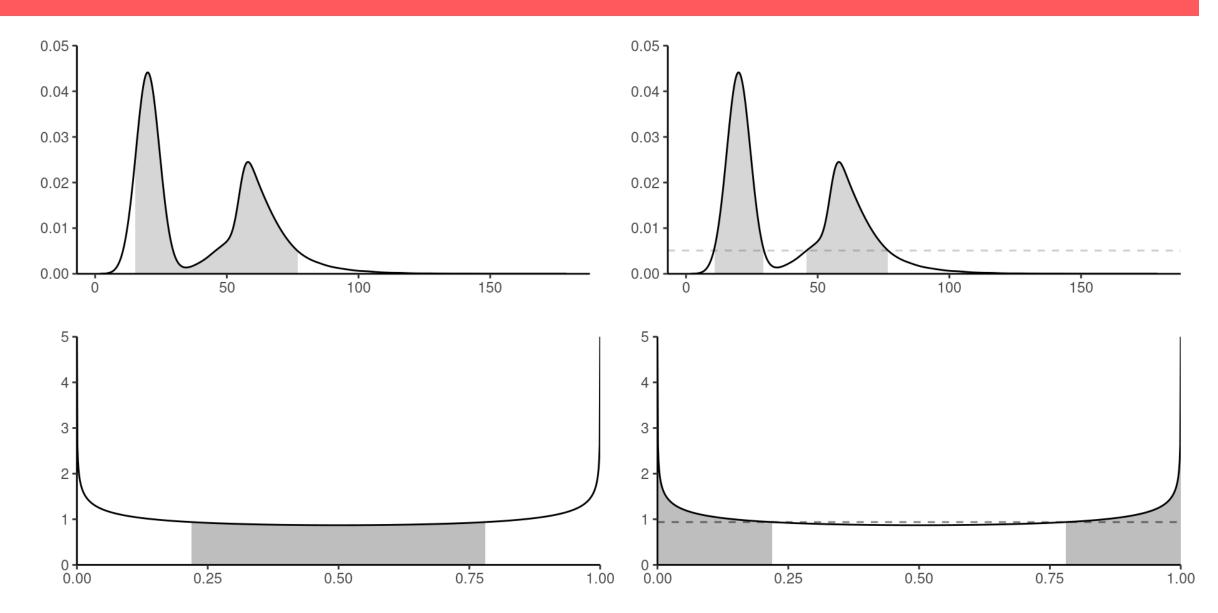


Figure 8: Density plots with 89% (top) and 50% (bottom) equitailed or central credible (left) and highest posterior density (right) regions for two data sets, highlighted in grey.

Computations of credible intervals

```
1 set.seed(2023)
 2 postsamp <- rbeta(n = 1000, shape1 = 0.5, shape2 = 0.2)
 3 alpha <- 0.11
 4 # Compute equitailed interval bounds
 5 quantile(postsamp, probs = c(alpha/2, 1-alpha/2))
     5.5%
              94.5%
0.0246807 0.9999980
 1 # Analytical (true) values
 2 qbeta(c(alpha/2, 1-alpha/2), shape1 = 0.5, shape2 = 0.2)
[1] 0.02925205 0.99999844
 1 # Highest posterior density intervals - note values are outside of the support!
 2 (hdiD <- HDInterval::hdi(density(postsamp), credMass = 1-alpha, allowSplit = TRUE))</pre>
           begin
                       end
[1,] -0.04331573 0.2800577
[2,] 0.47816030 1.1423868
attr(,"credMass")
[1] 0.89
attr(,"height")
[1] 0.3898784
```

References

Casella, G., & Berger, R. L. (2002). Statistical inference (2nd ed.). Duxbury.

Finetti, B. de. (1974). Theory of probability: A critical introductory treatment (Vol. 1). Wiley.

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